

Figure 1

MTIGEMLR SFLTKLEWFSTLFPRI PVPVQKNIDQQIKTRPRKI
KKDGKEGAEEIDRHVERRRSRSPRRSLSPRRSPRRSRSSH
REGHGSSSFDRELEREKERQRLEREAKEREKERRRSRSIDRG
LERRRSRSRERHRSRSRSRDRKGDRDRDREREKENERGR
RDRDYDKERGNEREKERERSRERSKEQSRGEVEEKKHKED
KDDRRHRDDKRDSKKEKKHSRSRSRERKHRSRSRSRNAGKR
SRSRSKEKSSKHKNESKEKSNKRSRSGSQGR TDSVEKSKKR
EHSPSKEKSRKRSRSKERSHKRDHSDSKDQSDKHDRRRSQSI
EQESQEKQHKNKDETV Stop

Figure 2

ACGCGGGGTTTCCAATTATTTGTTTCATTTATTTATTTTCTACATAACTAAATTAGAAACCTCACTGCTTCAT
 GGCAGTTGGTTTGCTATTGCTTCCAGTTTTATTAGGGCTTCATTTTATATTAGAGCTGTTAAAAGATAACCT
 TTAGACAGGAATTATCTAAAGTAGACATTTTATATTAGAGCTGTTAAAAGATAACCTTTAGACAGGAATTAT
 CTAAAGTAGATCATATGTAGCTAGGTTATGGTGCAAGGTGTATGATGTGTGCAAATATGTCCACAGAAATAA
 ATACATAGTAGGTATGTGGAATGTAAATTTAAGTCAATCGTTCCGCATAGTTTAGAAATGTAAGGGGCTTTT
 TCATATTGTTAACTGAGTGAGATCAGTTCCCTTTATGCCTGTGAGGCTGCAGGGTTTGTTCCTCACTTGCATG
 CACACACTAAGCCCAAATATTTCTGTTTCATTTCATTGTCAGATCAGGATATGAAAATAAAATTTTCTGTTAG
 TTTTTTTTGTATTGAGATCCCAAAGATGGTAATATTTTATAATATTCATGTATATATGGAAATACTTTTTT
 TGACGGCTAGGGTATCTTTTGTGTTTCTGTAGGACCTAGATGTGAAGGCTGGTGGAGGCTGTGTA
ATGACCATTGGAGAAATGCTACGATCTTTTCTCACAAAACCTGGAGTGGTTTTCTACCTTGTTTCCAAGAATT
 CCAGTTCCAGTTCAAAGAATATTGATCAACAGATTAAAACCCGACCTAGAAAAATCAAGAAAGATGGGAAG
 GAAGGTGCTGAGGAAATAGACAGACATGTTGAACGCAGACGTTCAAGGTCTCCAAGGAGATCTCTGAGTCCA
 CGGAGGTCCCCAAGAAGGTCAAGAAGTAGAAGTCATCATCGGGAGGGCCATGGGTCTTCTAGTTTTGACAGA
 GAATTAGAAAGAGAGAAAGAACGCCAGCGACTAGAGCGTGAAGCCAAAGAAAGGGAGAAAGAA
 CGGCGAAGATCCCCAAGTATTGACCGGGGGTTAGAACGCAGGCGCAGCAGAAGTAGGGAAAGGCATAGAAGT
 CGCAGTCGAAGTCGTGATAGGAAAGGGGATAGAAGGGACAGGGATCGAGAAAGAGAGAAAGAAAATGAGAGA
 GGTAGAAGACGAGATCGTGACTATGATAAGGAAAGAGGAAATGAACGAGAAAAAGAGAGAGAGCGATCAAGA
 GAAAGGTCCAAGGAACAGAGAAGTAGGGGAGAGGTAGAAGAGAAGAAACATAAAGAAGACAAGATGATAGG
 CGGCACAGAGATGACAAAAGAGATTCCAAGAAAGAGAAAAAACACAGTAGAAGCAGAAGCAGAGAAAGGAAA
 CACAGAAGTAGGAGTCGAAGTAGAAATGCAGGGAAACGAAGTAGAAGTAGAAGCAAAGAGAAATCAAGTAAA
 CATAAAAATGAAAGTAAAGAAAAATCAAATAAACGAAGTCGAAGTGGCAGTCAAGGAAGAACTGACAGTGTT
 GAAAAATCAAAAAAACGGGAACATAGTCCCAGCAAAGAAAAATCTAGAAAGCGTAGTAGAAGCAAAGAACGT
 TCCCACAAACGAGATCACAGTGATAGTAAGGACCAGTCAGACAAACATGATCGTCTGAAGGAGCCAAAGTATA
 GAACAAGAGAGCCCAAGAAAAACAGCATAAAAAACAAAGATGAGACTGTG
TGAAAATATTTTGTAAGAGTGGATCACATTGAATCCTATAAATGATTAAATCTGCTTTTTTCCCCACGTTG
 AGATTGTGCAGTAGTTCGCACTCCTCAAGCTCTCCCTGTAGGCTGCATTTTCATTTCTTCTTTCGTGTAGGG
 AAGTGCCTTTGTAATTCCATTTATTGTCATTGGTGTTCACCCAATTGTTAAGTTTGATACATGATGCACAG
 ATTGTTCTTGCATTTTTTATTGTTTGTTCCTGAAATGTACAGTCTGTACATATGTCCTGAAATGTTTTAATT
 CCTTTGGCATGGTTACCATGTTGGTTAAATTTGTATAAGGCAATAAACTGCCACTAATCCNAAAAAAAAAAAA
 AAAA~AAAA~AAAA~

Start codon (ATG) and stop codon (TGA) underlined

FIGURE 3

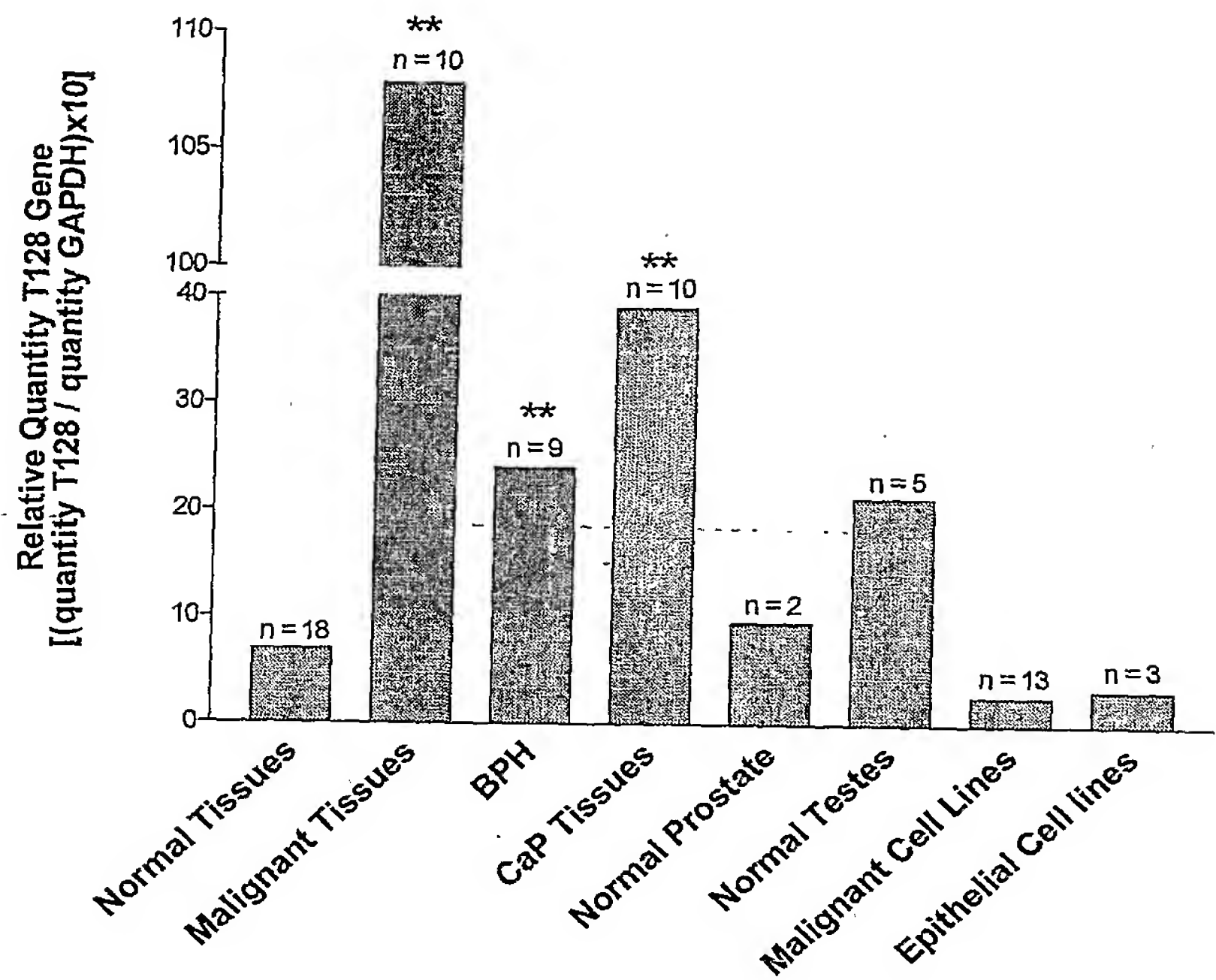


FIGURE 4

